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APPROVED O.G. FIG. ——
BY CLASS SUBCLASS
435 194
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GACCGCGCAGCTCAGCCTCCCGCCGATTGTATGTTCCAGGCCTCAATGAGGAGTCCAAA 60  $\mathbf{E}$ P F K Q Q K V E D F Y D Ι G 20 120 A I V C K K R E K S Т G 40 E Y GAGTGGCCAGTTTGCCATCGTGAAGAAGTGCCGGGAGAAGAGCACGGGGCTTGAGTATGC 180 F K K Ι R 0 S R S S R 60 E F AGCCAAGTTCATCAAGAAGCGGCAGAGCCGGGCGAGCCGGCGCGTGTGAGCCGGGAGGA 240 E S E V I L R Q V L Η Н N V I  $\mathbf{T}$ 80 Τ. Η GATCGAGCGGGAGGTGAGCATCCTGCGGCAGGTGCTGCACCACAATGTCATCACGCTGCA 300 V Y E N R  $\mathbf{T}$ D V L I v E L V S G G 100 CGACGTCTATGAGAACCGCACCGACGTGGTGCTCATCCTTGAGCTAGTGTCTGGAGGAGA 360 F D L Α 0 K E S L S E E E Α S F 120 420 Q I L D G V N Y L Н  $\mathbf{T}$ K I Η F D L 140 TAAGCAGATCCTGGATGGGGTGAACTACCTTCACACAAAGAAAATTGCTCACTTTGATCT 480 N T M L D K L N I ₽ I P H I K L I 160 CAAGCCAGAAAACATTATGTTGTTAGACAAGAATATTCCCATTCCACACATCAAGCTGAT 540 F G L Α E I H E D G V E F K N I 180 TGACTTTGGTCTGGCTCACGAAATAGAAGATGGAGTTGAATTTAAGAATATTTTTGGGAC 600 V Α P E I V N Y E P L G L E Α D 200 GCCGGAATTTGTTGCTCCAGAAATTGTGAACTACGAGCCCCTGGGTCTGGAGGCTGACAT 660 I G V I T Y T Τ. L S G G D 220 GTGGAGCATAGGCGTCATCACCTACATCCTCTTAAGTGGAGCATCCCCTTTCCTGGGAGA 720 Q E т L A Ι Т S N V S Y D F D E E F 240 CACGAAGCAGGAAACACTGGCAAATATCACATCAGTGAGTTACGACTTTGATGAGGAATT 780 S Η T S E L Α K D F I R K L V K Ε 260 CTTCAGCCATACGAGCGAGCTGGCCAAGGACTTTATTCGGAAGCTTCTGGTTAAAGAGAC 840 R Т Ι Q  $\mathbf{E}$ Α R L Η P W I  $\mathbf{T}$ P V D 280 CCGGAAACGGCTCACAATCCAAGAGGCTCTCAGACACCCCTGGATCACGCCGGTGGACAA 900 Q R R E Α M V S V V N L E N F 300 R K Q Y CCAGCAAGCCATGGTGCGCAGGGAGTCTGTGGTCAATCTGGAGAACTTCAGGAAGCAGTA 960 R R W K L S F S S I V L C N Η R L 320 TGTCCGCAGGCGTGGAAGCTTTCCTTCAGCATCGTGTCCCTGTGCAACCACCTCACCCG 1020 H R Ρ D E D L R C 340 E D CTCGCTGATGAAGAAGGTGCACCTGAGGCCCGGATGAGGACCTGAGGAACTGTGAGAGTGA 1080 Ι Α R R K Α L Н P R R R 360 CACTGAGGAGGACATcGCCAGGAGGAAAGCCCTCCACCCACGGAGGAGGAGCAGCACCTC 1140 CTAAcTGGCCTGACCTGCAGTGGCCGCCAGGGAGGTTTGGGCCCAGCGGGGCTCCCTTCT 1200 GTGCAGACTTTTGGACCCAGCTCAGCACCAGCACCCGGGCGTCCTGAGCACTTTGCAAGA 1260 GAGATGGGCCCAAGGAATTCAGAAGAGCTTGCAGGCAAGCCAGGAGACCCTGGGAGCTGT 1320 GGCTGTCTTCTGTGGAGGAGGCTCCAGCATTCCCAAAGCTCTTAATTCTCCATAAAATGG 1380 1440 ATAAAGGACATCCTCATCACGGGGTGAAGGTCAGAGTAAGGCAGCCTTCTTCACAGG 1500 CTGAGGGGGTTCAGAACCAGCCTGGCCAAAAATTACACCAGAGAGACAGAGTCCTCCCCA 1560  ${ t TTGGGAACAGGGTGATTGAGGAAAGTGAACCTTgGGTGTGAGGGACCAATCCTGTGACCT$ 1620 CCCAGAACCATGGAAGCCAGGACGTCAGGCTGaCCAACACCTCAGACCTTCTGAAGCAGC 1680 CCATTGcTGGCCCGCCATGTTGTAATTTTGCTCATTTTTATTAAActtctggtttacctg 1740 aa 1742

	F1G. 2A
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	13 YDTGEELGSGQFAVVKKCREKSTGLQYAAKFI 13 YEMGEELGSGQFAIVKKCROKGTGKEYAAKFI 13 YDIGEELGSGQFAIVKKCREKSTGLEYAAKFI 61 LCPGRELGRGKFAVVRKCIKKDSGKEFAAKFM 33 ILTSKELGRGKFAVVRQCISKSTGQEYAAKFL
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	46 KRRTKSSRRGVSREDTEREVSILKEI-QHPNVI 46 KRRLSSSRRGVSREETEREVNILREI-RHPNTI 46 KRQSRASRRGVSREETEREVSILRQV-LHHNVI 94 KRRKGQDCRMETIHETAVLELAQDNPWVI 66 KRRRGQDCRAETLHETAVLELAKSCPRVI
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	78 TLHEVYENKTDVILILELVAGGELFD-FLAEK- 78 TLHDIFENKTDVVLILELVSGGELFD-FLAEK- 78 TLHDVYENRTDVVLILELVSGGELFD-FLACK- 123 NLHEVYETASEMILVLEYAAGGEIFDQCVADRE 95 NLHEVYENTSEIILILEYAAGGEIFSLCLPELA
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	109 ESLTEEEATEFLKQILNGVYYLHSLQIAHFDLK 109 ESLTEDEATQFLKQILDGVHYLHSKRIAHFDLK 109 ESLSEEEATSFIKQILDGVNYLHTKKIAHFDLK 156 EAFKEKDVQRLMRQILEGVHFLHTRDVVHLDLK 128 EMVSENDVIRLIKQILEGVYYLHQNNIVHLDLK
DÄP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	PENIMILDRNVPKPRIKIIDFGLAHKIDFGNEF 142 PENIMLLDKNVPNPRIKLIDFGIAHKIEAGNEF 142 PENIMLLDKNIPIPHIKLIDFGLAHEIEDGVEF 189 PONILLTSESPLGD-IKIVDFGLSRILKNSEEL 161 PONILLSSIYPLGD-IKIVDFGMSRKIGHACEL
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	175 KNIFGTPEFVAPEIVNYEPLGLEADMWSIGVIT 175 KNIFGTPEFVAPEIVNYEPLGLEADMWSIGVIT 175 KNIFGTPEFVAPEIVNYEPLGLEADMWSIGVIT 221 REIMGTPEYVAPEILSYDPISMATDMWSIGVLT 193 REIMGTPEYLAPEILNYDPITTATDMWNIGIIA
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	208 YILLSGASPFLGDTKQETLAMVSAVMYEFEDBY 208 YILLSGASPFLGETKQETLTNISAVMYDFDEEY 208 YILLSGASPFLGDTKQETLANITSVSYDFDEEF 254 YVMLTGISPFLGNDKQETFLNISQMNLSYSEEE 226 YMLLTHTSPFVGEDNQETYLNISQVNVDYSEET
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	FSNTSALAKDFIRRLLVKDPKKRMTIQDSLQHP FSNTSELAKDFIRRLLVKDPKRRMTIAQSLEHS FSHTSELAKDFIRKLLVKETRKRLTIQEALRHP FDVLSESAVDFIRTLLVKKPEDRATAEECLKHP FSSVSQLATDFIQSLLVKNPEKRPTAEICLSHS
DAP-kinase ZIP-kinase DRP-1 DRAK1 DRAK2	274 WI 274 WI 274 WI 320 WL 292 WL

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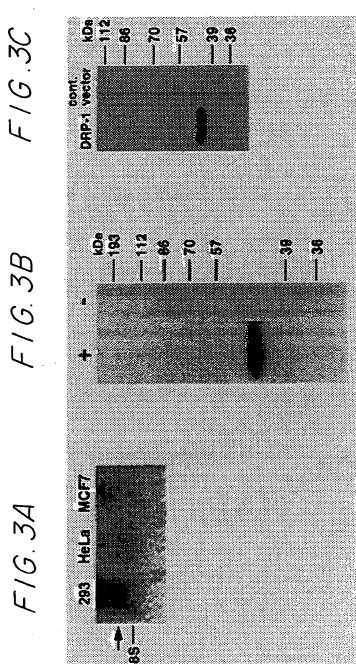
0.G. FIG. —	CLASS SUBCLASS	435 144	
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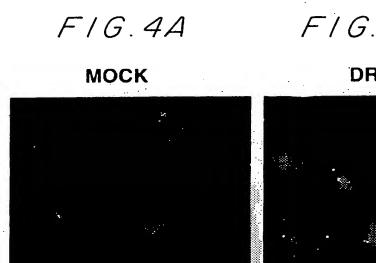
## F16.2B

DAP-kinase	1	MMEKFKKFAARKKWKDSVRLISLCORLSR	29
DRP-1	1	NLENFIKQYVRRRWKLSFSIVSLCNHLTR	29
smMLCK	1	SKORMKKYMARRKWOKTGHAVRAIGRLSS	29
CaMKIIa	1	TVDCLKKLNARRKLKGAILTTMLATRNFS	29
CaMKI	1	VSEQIKKNFAKSKWKDAFNAT-AVVRHMR	28
CaMKIV	1	MDTAOKKLOEFNARRKIKAAVKAVVASSRLGS	32
ZIP-kinase	1	GEDSGRKPERRIKTTRLKEYTIKSHSS	28

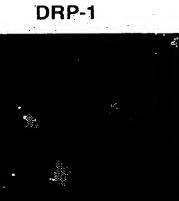
APPROVED O.G. FIG

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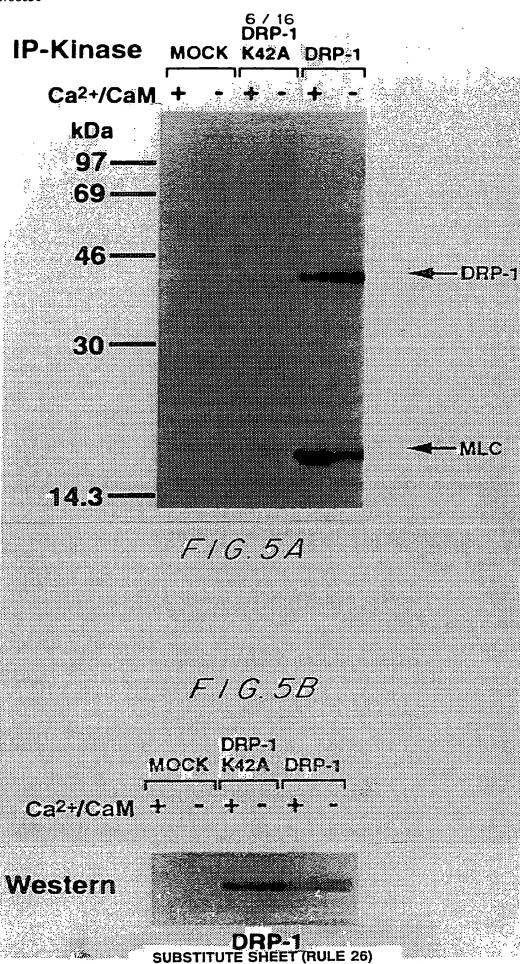


F1G.4B



F1G.4C

**DAP-kinase** kDa 193 112 86



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F16.6B

F16.60

F16.64

F16.6C

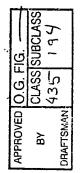


FIG.7

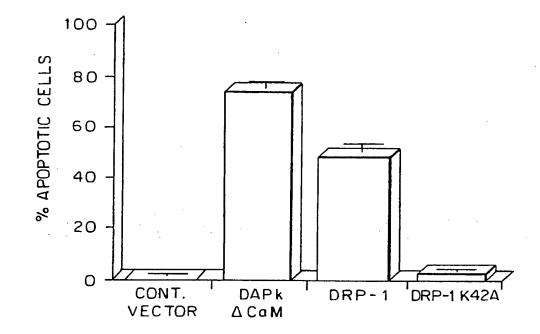
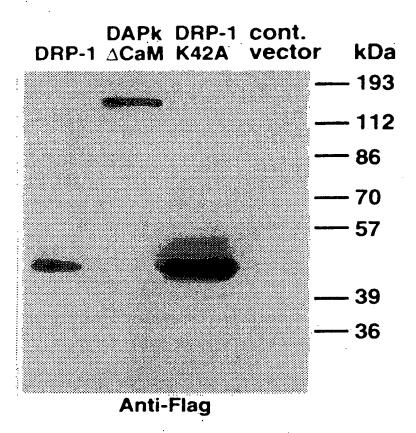
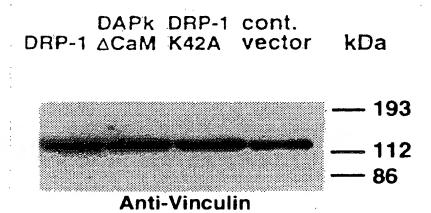


FIG.8A



F1G.8B



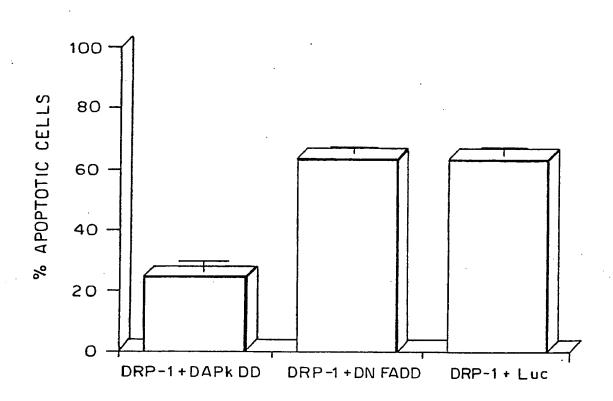
APPROVED O.G. FIG.

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FIG. 9A





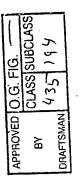
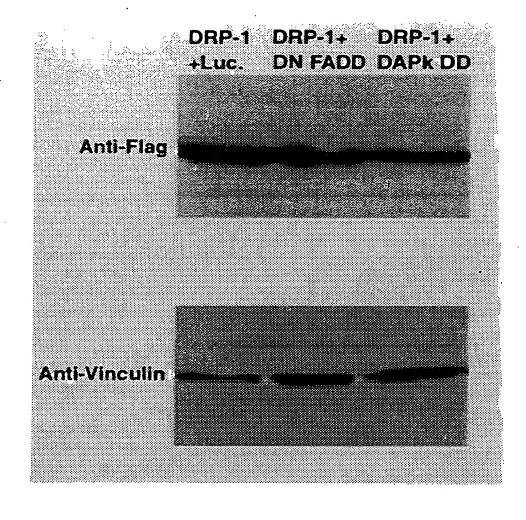


FIG 9B





F1G.10A

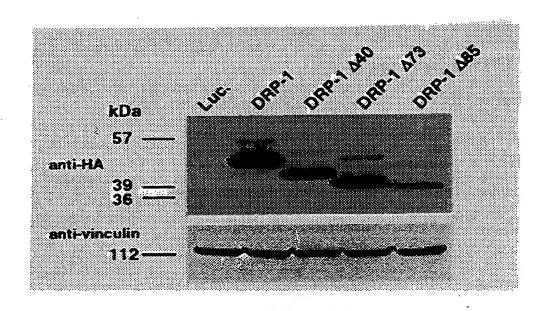
KINASE CaM Reg.
DOMAIN
DRP-1 1 1 360

DRP-1 Δ40 1 1 320

DRP-1 Δ73 1 1 287

DRP-1 Δ85 1 1 275

F1G.10B



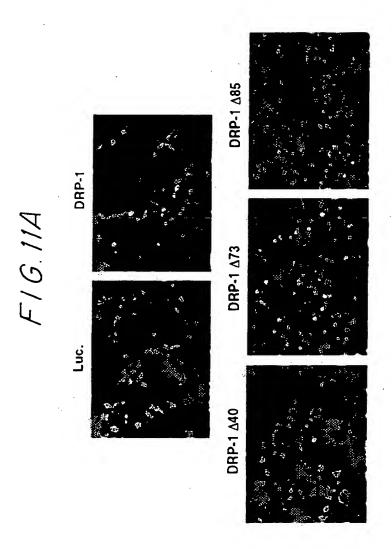
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APPROVED O.G. FIG.

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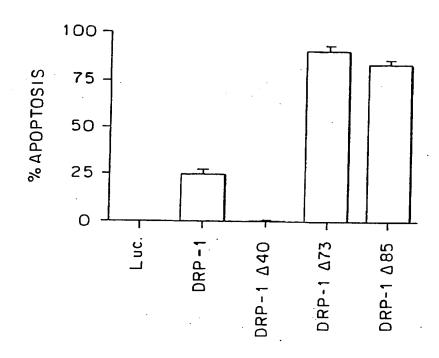
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,	0.G. FIG. —	CLASS SUBCLASS	435 147
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F1G. 11B



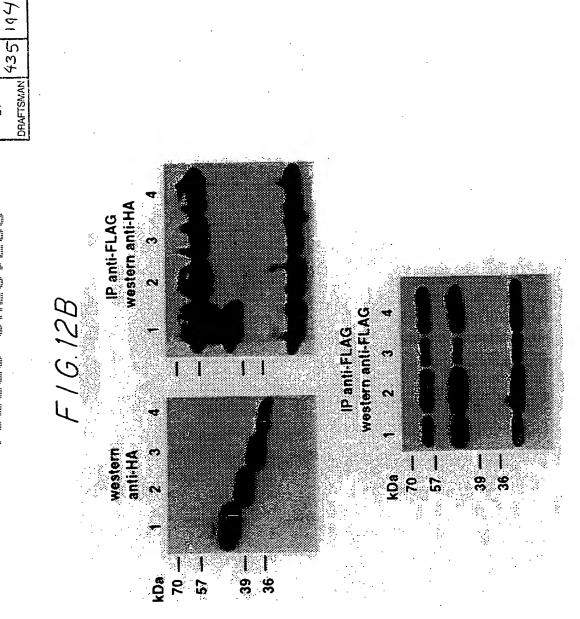
O.G. FIG. —	CLASS SUBCLASS	432 144	
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O.G. FIG. —	CLASS SUBCLASS	435 194
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	45a 193 — 112 — 86 —	70 <del>-</del> 57 <del>-</del>	36 —	
FLAG anti-HA				
IP anti-FLAG western anti-HA	`  - 			
western anti-FLAG	1			
	KDa 193 — 112 — 86 —	70 — 57 —	38 —	

APPROVED O.G. FIG.

β



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